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FIG. 1A

SEQ ID NO: 1	MtubZ94752	<u>mlrlvvqalllvlafaggyavaacktvltltvdgtamrvttmkrsvidive</u>	50
	MtubZ94752	engfsvddrddlypaagvqvhdadtivlrrsrplqisldghdkgvwtta	100
	MtubZ94752	stvdealaqlamtdtapaaasrasrvplsgmalpvvsaktvqlndgglvr	150
SEQ ID NO: 2	MtubZ94752	tvhlpapnvagllsaagvp11qsdhvvpaatapivegmgiqvtrnrkkr	200
	MtubMTV008	----- <u>mpvqwlwrartakgttlknarttlaaaiaqt</u>	32
SEQ ID NO: 3	Mlep104666	-----mseyrkl	8
SEQ ID NO: 4	MtubMTV043	-----msgrhrkpt	9
	MtubZ94752	terlplppnarrvedpemmsrevvedpgvpqtdvtfavaevngvetgr	250
SEQ ID NO: 36	MlutZ96935	-----mtlfttsat	9
SEQ ID NO: 5	MlepL01095	-----mpqemldvrklc	12
SEQ ID NO: 6	MtubU38939	-----mhplpadhgrsrcnrhplspisplsignisatsqdmssmt	38
SEQ ID NO: 7	MtubZ81368	-----mtpqllttagaqrprdrca	19
	MtubMTV008	<u>lvttspagianaddagldpnaaagpdavgfdpnlppapdaapvdtppape</u>	82
SEQ ID NO: 8	Scoeli6C12\$	---irtaavtlvaatalgatgeavaapsaplrtDWDAIAACESSGNWQAN	25
	Mlep104666	ttssiiivakitftgamldqsgialagqaspatdsEWDQVARGESGGNWSIN	58
	MtubMTV043	tsnvsvakiaftgavlgqgggiamaaqataatdgEWDQVARGESGGNWSIN	59
	MtubZ94752	lpvanvvtpaheavrvrgtkpgtevppvidgsIWDIAIGCEAGGNWAIN	300
	MlutZ96935	rsrratasivagmtlagaaavqfsapagaatvdTWDRLAECESNGTWDIN	59
	MlepL01095	klfvksavvsqivtasmlststgmanavprePNWDAVAQCESGRNWRAN	62
	MtubU38939	riakpliksamaaglvtasmslstavahagpsPNWDAVAQCESGGNWAAN	88
	MtubZ81368	rivctvfietavvatmfvallqlstisskaddIDWDAIAQCESGGNWAAN	69
	MtubMTV008	dagfdpnlppplapdfllspaaeeappvpvaysVNWDAIAQCESGGNWSIN	132
		.** *	
	Scoeli6C12\$	TGNGYYGGLQFARSSWIAAGGLKYAPRADLATRGEQIAVAERLARLQGMS	75
	Mlep104666	TGNGYLGGLOFSQGTWASHGGGEYAPSAQLATREQQIAVAERVLATQSGG	108
	MtubMTV043	TGNGYLGGLOFTQSTWAAHGGGEFAPSAQLASREQQIAVGERVLATQGRG	109
	MtubZ94752	TGNGYYGGVQFDQGTWEANGGLRYAPRADLATREEQIAVAEVLRLRQGWG	350
	MlutZ96935	TGNGFYGGVQFTLSSWQAVGGEG---YPHQASKAEQIKRAEILQDLQGWG	106
	MlepL01095	TGNGFYGGLOFKPTIWARYGGVG---NPAGASREQQITVANRVLADQGLD	109
	MtubU38939	TGNGKYGGLOFKPATWAAFGGVG---NPAAASREQQIAVANRVLAEQGLD	135
	MtubZ81368	TGNGLYGGLOISQATWDSNGGVG---SPAAASPOQQIEVADNIMKTQPGG	116
	MtubMTV008	TGNGYYGGLQFTAGTWRANGGSG---SAANASREEQIRVAENVLRSQGIR	179
		****.***** . * . ** * .***** *** .. **	
	Scoeli6C12\$	AW-----	78
	Mlep104666	AWPACGHGLSGPSLQEVLPAG---MGAPw----INGAPAPLAPPPPAEPAP	152
	MtubMTV043	AWPVCGRGLSNATPREVLPA SaamDAPldaaaVNGEPAPLA-PPPADPAP	158
	MtubZ94752	AWPVCAaragar-----	362
	MlutZ96935	AWPLCSQKLgltqadadagdvdateaapvavertatvqrqsaadeaaaeg	156
	MlepL01095	AWPKCGAASDLPLITLWSHPAQGVKQIINDIIImgdttlaaialngl----	155
	MtubU38939	AWPTCGAASGLPIALWSKPAQGIKQIINEIIiwagiqasipr-----	176
	MtubZ81368	AWPKCSscsqgdaplgslthiltflaaetggcsgsrdd-----	154
	MtubMTV008	AWPVCGrng-----	188
		*** *	

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FIG. 1A (CONT.)

Mlep104666	pqppadnf-----PPTPGDVPSPLarp-----	174
MtubMTV043	pvelaandlpaplgelplpaapadpappadlaPPAPADVAPPVelavndlp	208
MlutZ96935	aaaaeqavvaeaetivvksgdslwtlaneyeveggwtalyeankgavsda	206
MtubMTV043	aplgelplpaapadpappadlappapadlappapadlappapadlappvel	258
MlutZ96935	aviyvgqelvlpqa-----	220
MtubMTV043	avndlpaplgelplpaapaelappadlapasadlappapadlappapaela	308
MtubMTV043	ppapadlappaavneqtapgdqpatapggpvglatdlelpepdppadap	358
MtubMTV043	ppgdvteapaetpqvsniaytkklwqairaqdvcgndaldslagpyvig-	407

Fig. 1B

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[illegible]

SEQ ID NO:	RPF	TIYVKS	GD	SL	WT	LANE	YE	VE	EG	WT	AL	YE	ANK	GA	VS	-----	DA	VI	YV	GQ	EL	VL	PP	QA	
13	g149657	TIYVKS	GD	SL	WT	LANE	YE	VE	EG	WT	AL	YE	ANK	GA	VS	-----	DA	VI	YV	GQ <td>EL</td> <td>VL</td> <td>PP</td> <td>QA</td>	EL	VL	PP	QA	
14	g2226145	TIYVKS	GD	SL	WK	LS	RQ	YD	IT	--	IS	AL	KS	EN	KL	-----	KS	TV	LY	VG	QS	SL	KV	PS	
15	g2226145	TIYVKS	GD	SL	WK	LA	QT	YN	TS	--	VA	AL	TS	AN	IL	-----	ST	TV	LS	IG	QT	LT	IP	--	
16	g2226145	TYTVKS	GD	SL	WV	IA	QK	FN	VT	--	AQ	IR	EK	NN	-----	KT	DV	LQ	VG	QK	LV	-----	-----		
17	g2226145	KYTVKS	GD	SL	WK	IA	NN	IN	LT	--	VQ	IR	NI	NN	-----	KS	DV	LY	VG	QV	LV	KL	-----	-----	
18	g266725	TYTVKS	GD	TI	WAL	SS	K	YG	TS	--	VQ	NI	MS	WN	NN	-----	SS	SS	IV	VG	QV	LV	AV	KQ	
19	g80581	THAVKS	GD	TI	WAL	SV	K	YG	VS	--	VQ	DI	MS	WN	NN	-----	SS	SS	IV	VG	QK	LA	IK	Q	
20	g2707292	SVVKKS	GD	TI	WAL	SV	K	YK	TS	--	IA	Q	L	KS	WN	NH	-----	SS	DT	IY	IG	QN	LI	VS	QS
21	g755216	TYTVKS	GD	TI	WG	IS	QR	Y	GIS	--	VA	Q	I	Q	S	AN	NN	-----	KS	T	IY	IG	Q	L	LL
22	g1722873	TYTVKK	GD	TI	LD	IA	GR	FY	GN	ST	QW	RK	IWN	ANK	TAM	IK	RS	KRN	IR	Q	P	GH	WI	FP	GQ
23	g1176755	TYTVKK	GD	TI	LD	IA	GR	FY	GD	ST	QW	RK	IWN	VKK	KAM	IK	RS	KRN	IR	Q	P	GH	WI	FP	GQ

FIG. 1C

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SEQ ID NO: 4 1 msgrhrkpttsnvsvakiaftgavlrggggiamaaqataatdgewdqvarcesgggnwsintgngylgg
 lqftqstwaahggggefapsaqlasreqqiavgervlatqgrgawpvcgrglstnatprevlpasaamd
 apldaaavngepaplapppadv 156

157 appvelaandlpaplgelplaapadpappadlappapadv 196
 197 appvelavndlpaplgelplaapadpappadlappapadlappapadl 252
 253 appvelavndlpaplgelplaapaelappadlap-asadlappapadlappapaelappapadlappa
 320 -----avne 323

324 qtapgdqpatapggpvglatdlepdpqpapppgdvteapaetpqvsniaytkklwqaira
 389 qdvcgndaldslagpyvig* 407

Motif	sequence
A	157 appvelaandl 167 SEQ ID NO: 25
B'	168 paplgelplaapad 181 SEQ ID NO: 28
C	182 pappadl 188 SEQ ID NO: 29
D	189 appapadv 196 SEQ ID NO: 31
A	197 appvelavndl 207 SEQ ID NO: 26
B'	208 paplgelplaapad 221 SEQ ID NO: 28
C	222 pappadl 228 SEQ ID NO: 29
D	229 appapadl 236 SEQ ID NO: 30
D	237 appapadl 244 SEQ ID NO: 30
D	245 appapadl 252 SEQ ID NO: 30
A	253 appvelavndl 263 SEQ ID NO: 26
B	264 paplgelplaapael 278 SEQ ID NO: 27
C	279 appadl 284 SEQ ID NO: 55
D*	285 apasadl 291 SEQ ID NO: 56
D	292 appapadl 299 SEQ ID NO: 30
D	300 appapael 307 SEQ ID NO: 57
D	308 appapadl 315 SEQ ID NO: 30
D'	316 appa 319 SEQ ID NO: 58
'A'	320 avne 323 SEQ ID NO: 59

A = appvela[av]ndl

B = paplgelplaapa[de]l

C = pappadl

D = appapa[de][lv]

FIG. 1D

SEQ ID NO:34 Lmonocytoγ.. 72
SEQ ID NO: 36 MlutFactor 62

nnkkatlaaaglavafaplaasastvvveagdtlwgiaqskgttvdaikkannlttqkivpggkigv
ttfttatrtrratagivagmllagaaagfsapaqaat-----vdtwrlaecaesngtwdintg
nevaaaekpekvsatwlnvrtgagvndnlltsaggtkvvtettesngwhkityndgktgfvngkyltdka
gfyggvffllswwqavgggyphq---aakaegrraeillqdlqgwawplcsqkigltqadaaag-----

144
125

Lmonocytoγ..
MlutFactor

vstpvpaptqevkketttttqqnnevaetktvkvktttttpppkvvaetkettvtdqonettthavksgbntiivansv
vvdatttannpvaiveratvtrqrsadedaeeaaeqaaaadqavvaeetivksgdislvttan

216
184

Lmonocytoγ..
MlutFactor

ktgsvvdimswnnl-----sssslavvgoekdaikttantatpkavkteapaaekqaapvvkentntntatt
evveggwtalyeankgavsdAAVlavvgoevlvpaa-----

283
220

Lmonocytoγ..
MlutFactor

ekketatqqqtapkapteaakpapapstntnankntntntntntpskntntnsntntnsntnanggss-----

355
220

Lmonocytoγ..
MlutFactor

nnnsnssasaliaeaqkhlkgayswgngpttfdcsgytkyvfakagislprtsgaqyasttrisesqakpg-----

427
220

Lmonocytoγ..
MlutFactor

dlvffdygsgishvglyvngngminaqdngvkydnhgsgwgkylvgfgrv-----

478
220

FIG. 1E

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SEQ ID NO: 35 1 accaaggagaaggacgaccccggtgtgcctcggccgcccgatcagcgaggactcgccatgg 60
 61 acaccatgactctcttcaccacttccgccacccgctcccgcggtgccaccgcctcgatcg 120
 M T L F T T S A T R S R R A T A S I V
 SEQ ID NO: 37 g
 121 tcgcgggcatgaccctcgccggcgccgcccgtgggcttctccgccccggcccaggccg 180
 A G M T L A G A A A V G F S A P A Q A A
 SEQ ID NO: 38 A
 oligo A1>>>
 csacsgtsgacacstgggaccgscstsgcsgag
 181 ccaccgtggacacctgggaccgcctcgccgagtgcgagtgccaacggcacctgggacatca 240
 T V D T W D R L A E C E S N G T W D I N
 T V D T W D R L A E E x S N G T x D
 <<< oligo G2 SEQ ID NO: 40 oligo G1>>>
 gttgccgaagatgccgcc agttcaccctgtcctcctg
 241 acaccggcaacggcttctacggcgccggtgcagttcaccctgtcctcctggcaggccgctcg 300
 T G N G F Y G G V Q F T L S S W Q A V G
 SEQ ID NO: 42 G
 <<< oligo A2
 SEQ ID NO: 41 ccictycciatrggigtrgtycg
 301 gcggcgaaggctacccgcaccaggcctcgaaggccgagcagatcaagcgcgccgagatcc 360
 G E G Y P H Q A S K A E Q I K R A E I L
 G E G Y P H Q A S K
 361 tccaggacctgcagggctggggcgcggtggccgctgtgctcgcagaagctgggcctgaccc 420
 Q D L Q G W G A W P L C S Q K L G L T Q
 421 aggctgacgcggacgcccgtgacgtggacgccaccgaggccgccccggtcgccgtggagc 480
 A D A D A G D V D A T E A A P V A V E R
 481 gcacggccaccgtgcagcgccagtcgcccgaggctgccgcccagcaggccgctg 540
 T A T V Q R Q S A A D E A A A E Q A A A
 541 ccgcgagcagggcgtcgtcgccgaggccgagaccatcgtcgtcaagtccggtgactccc 600
 A E Q A V V A E A E T I V V K S G D S L
 601 tctggacgctcgccaacgagtacgaggtggaggggtggctggaccgccctctacgaggcca 660
 W T L A N E Y E V E G G W T A L Y E A N
 661 acaaggcgccgtctccgacgcccgtgatctacgtcgccaggagctcgtcctgcccgc 720
 K G A V S D A A V I Y V G Q E L V L P Q
 721 aggcctgagacgcctgaccggccccccggaccggtacc 758
 A *

SEQ ID NO: 43 1 ATVDTWDRLE ECESNGTWDI NTGNIFYGGV QFTLSSWQAV GGEGYPHQAS KAEQIKRAEI 60
 61 LQDLQGWGAW PLCSQKLGLT QADADAGDVD ATEAAPVAVE RTATVQRQSA ADEAAAEQAA 120
 121 AAEQAVVAEA ETIVVKSGDS LWTLANEYEV EGGWTALYEA NKGAVSDAAV IYVQELVLP QA 182

FIG. 2A

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SEQ ID NO: 44 ggatccgcaccgccgcggttaaccctggtcgccgcgaccgcactcggggcgaccggcgaag 60
SEQ ID NO: 45 I R T A A V T L V A A T A L G A T G E A
cggtggccgcgccctcggcgcccctgcgcaccgactgggacgccatcgccgcgtgcgagt 120
V A A P S A P L R T D W D A I A A C E S
ccagcggcaactggcaggcgaacaccggcaacggctactacggcggcctgcagttcgcac 180
S G N W Q A N T G N G Y Y G G L Q F A R
gggccagctggatcgccgccggcgccctcaagtaacgccccgcgcgcggacctcgccaccc 240
S S W I A A G G L K Y A P R A D L A T R
gcggcgagcagatcgccgtggcggaacgcctcgcccgtctgcaggggatgtccgcctgg 299
G E Q I A V A E R L A R L Q G M S A W

FIG. 2B

09445289-054100

FIG. 3

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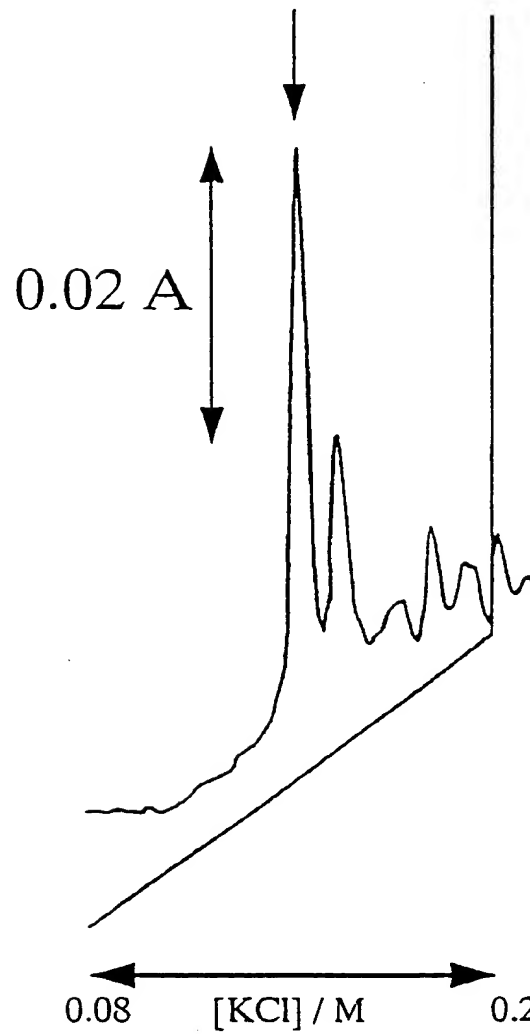
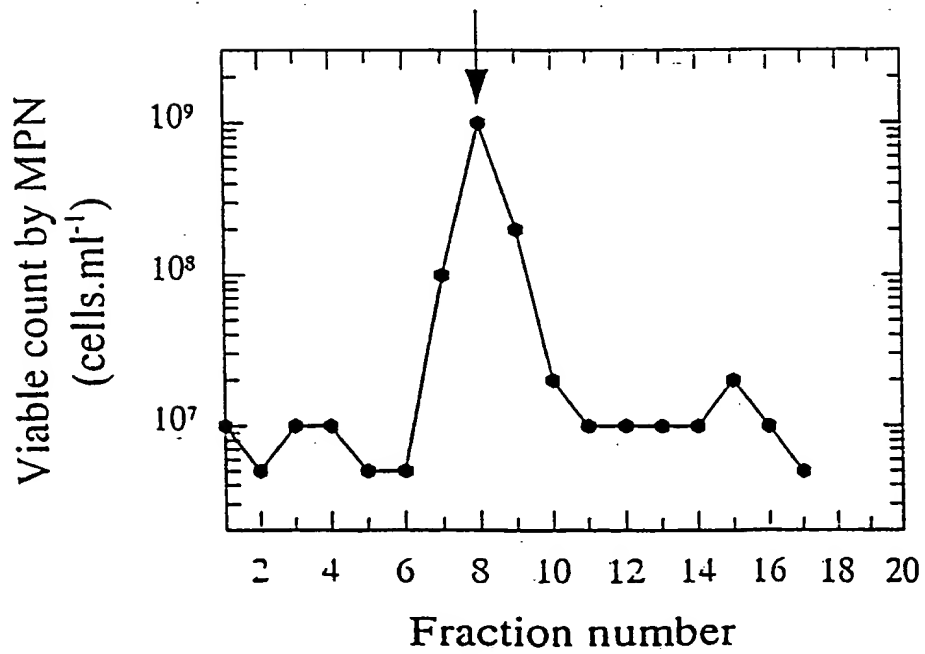
A**B**

FIG. 3

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C.

 M_r (kDal)94
66

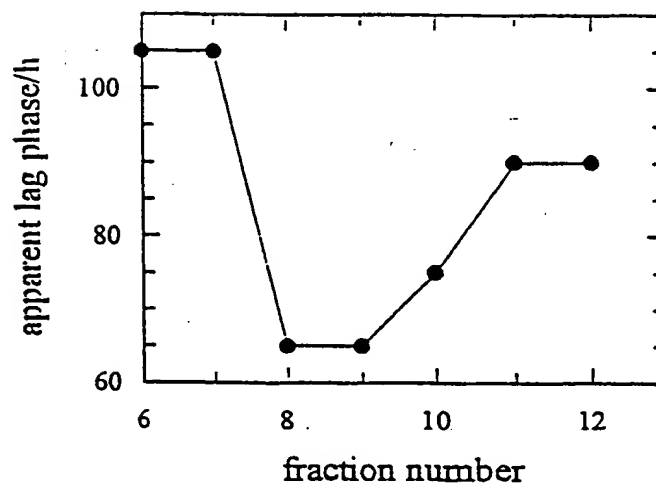
20.1

14.4

1

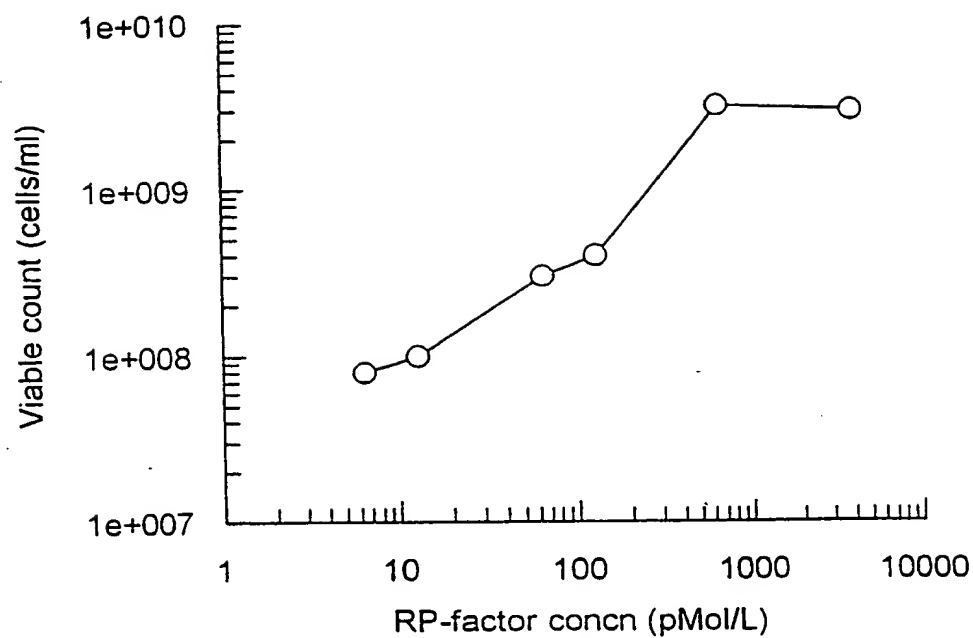
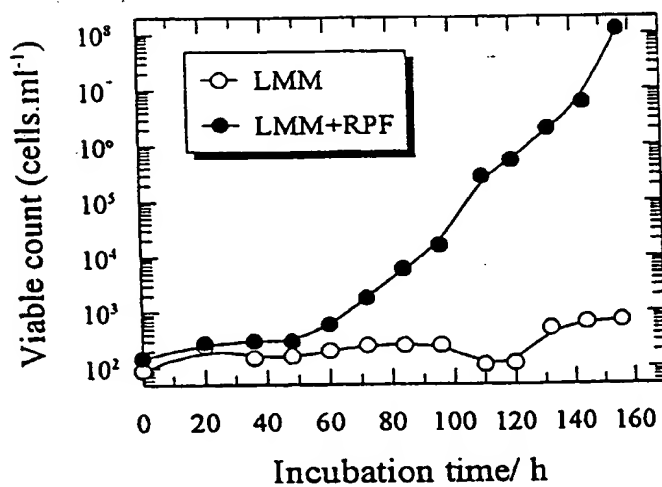
2

3



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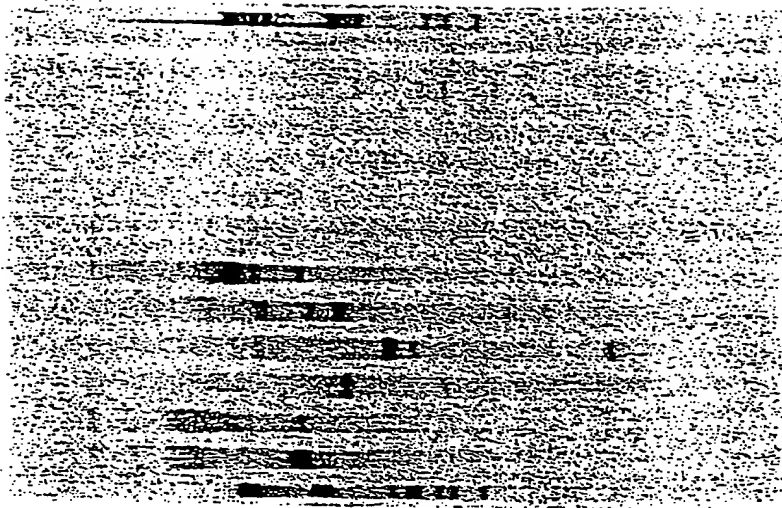
FIG. 4

A**B**

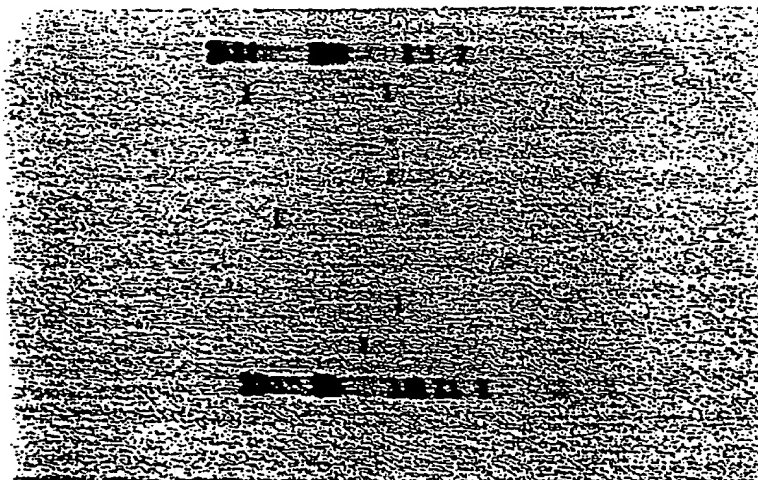
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FIG. 5

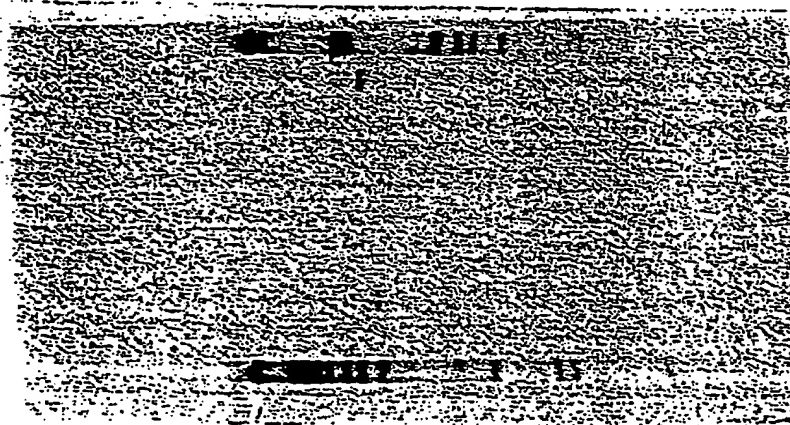
C



B



A



000000 63254450

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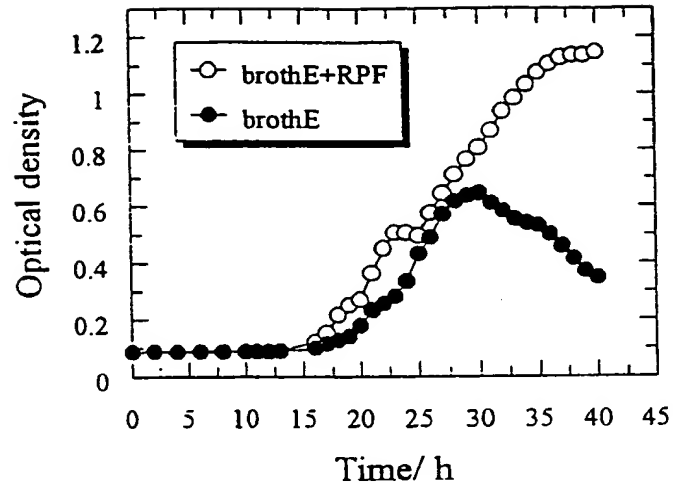
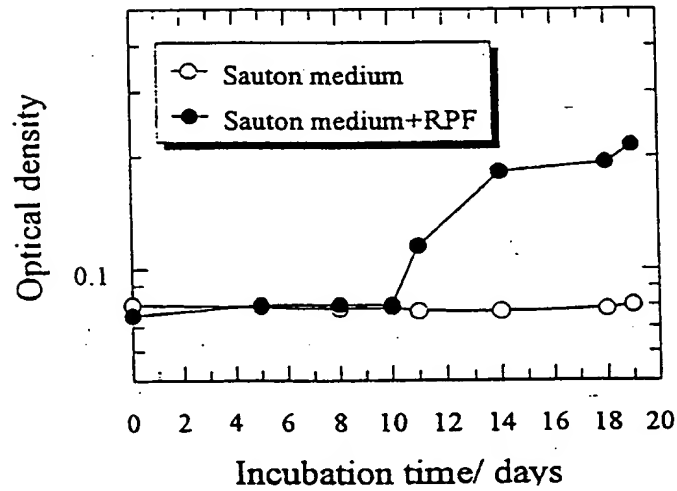
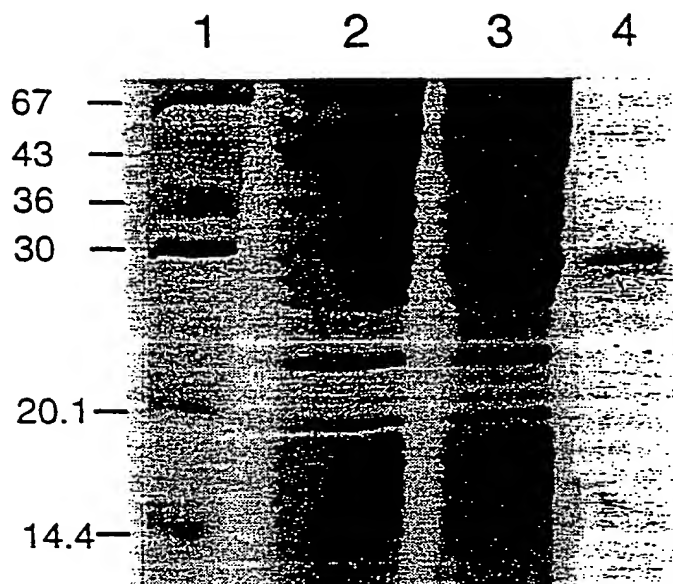
A**B**

FIG. 6

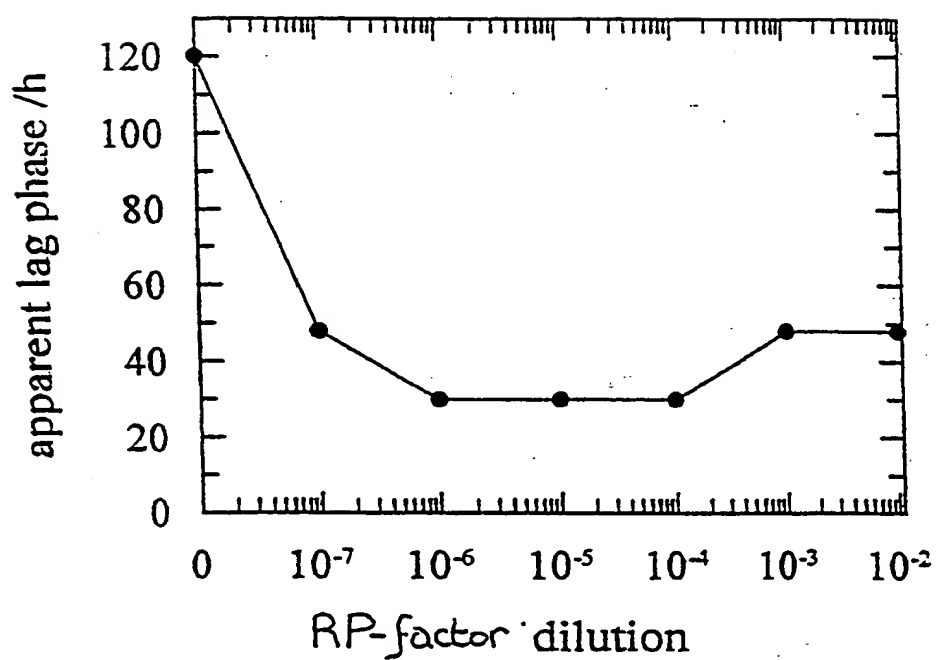
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FIG. 7

A



B



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C

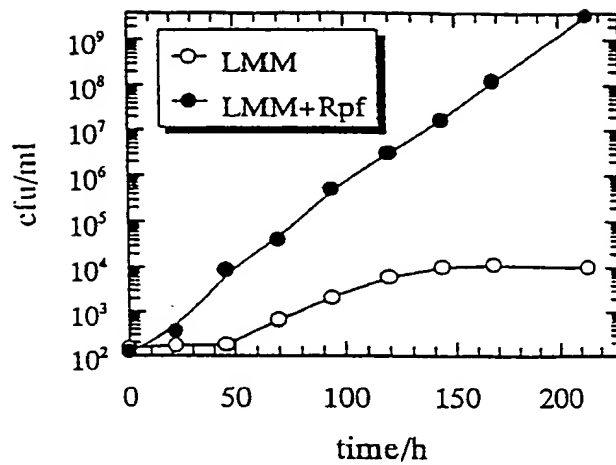


FIG. 7

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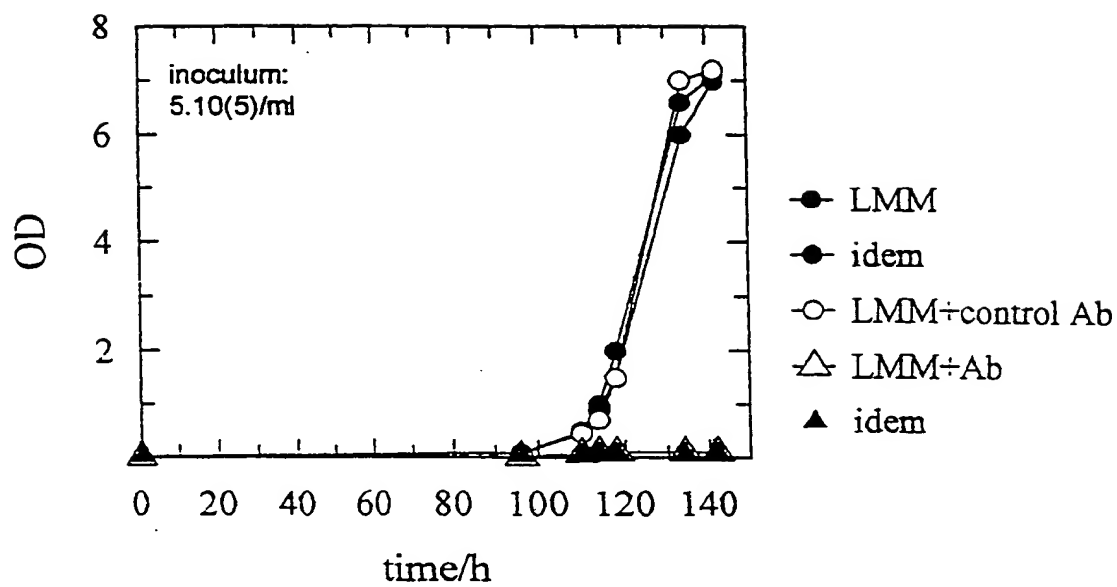
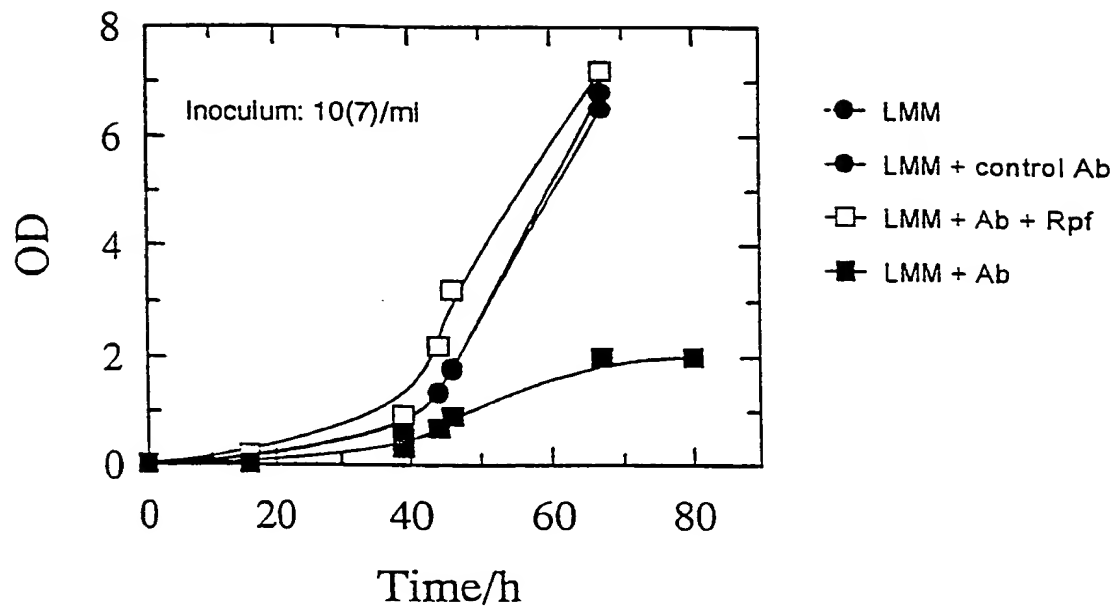


FIG. 8A

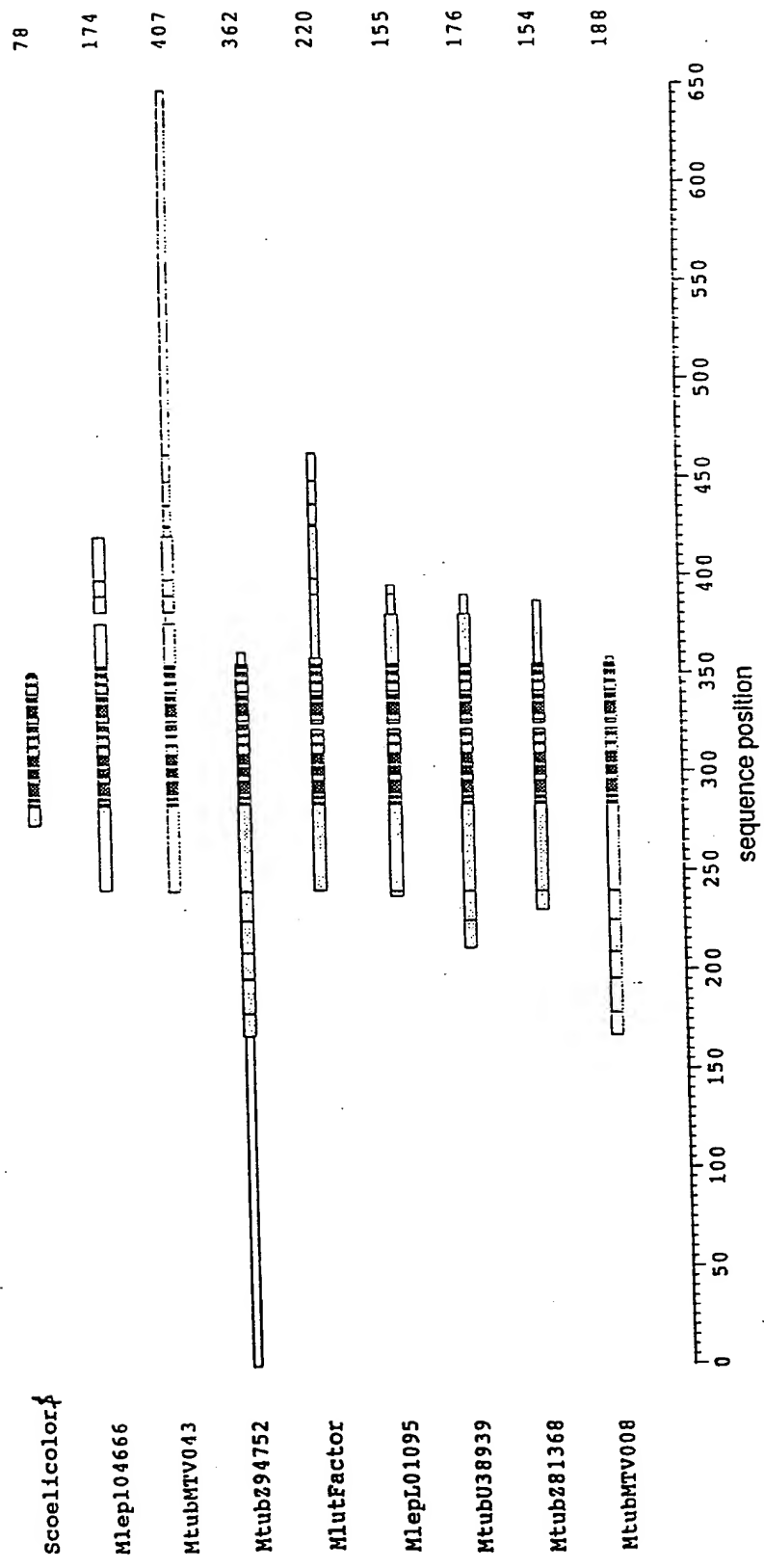
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FIG. 8 B



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FIG. 9A



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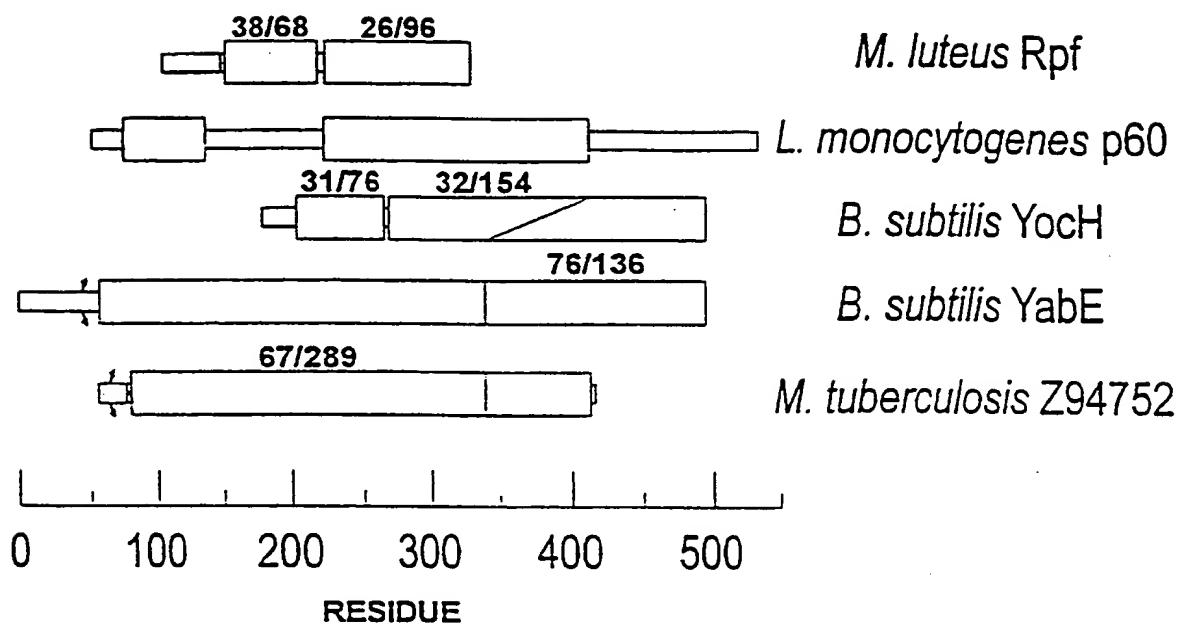


FIG. 9B

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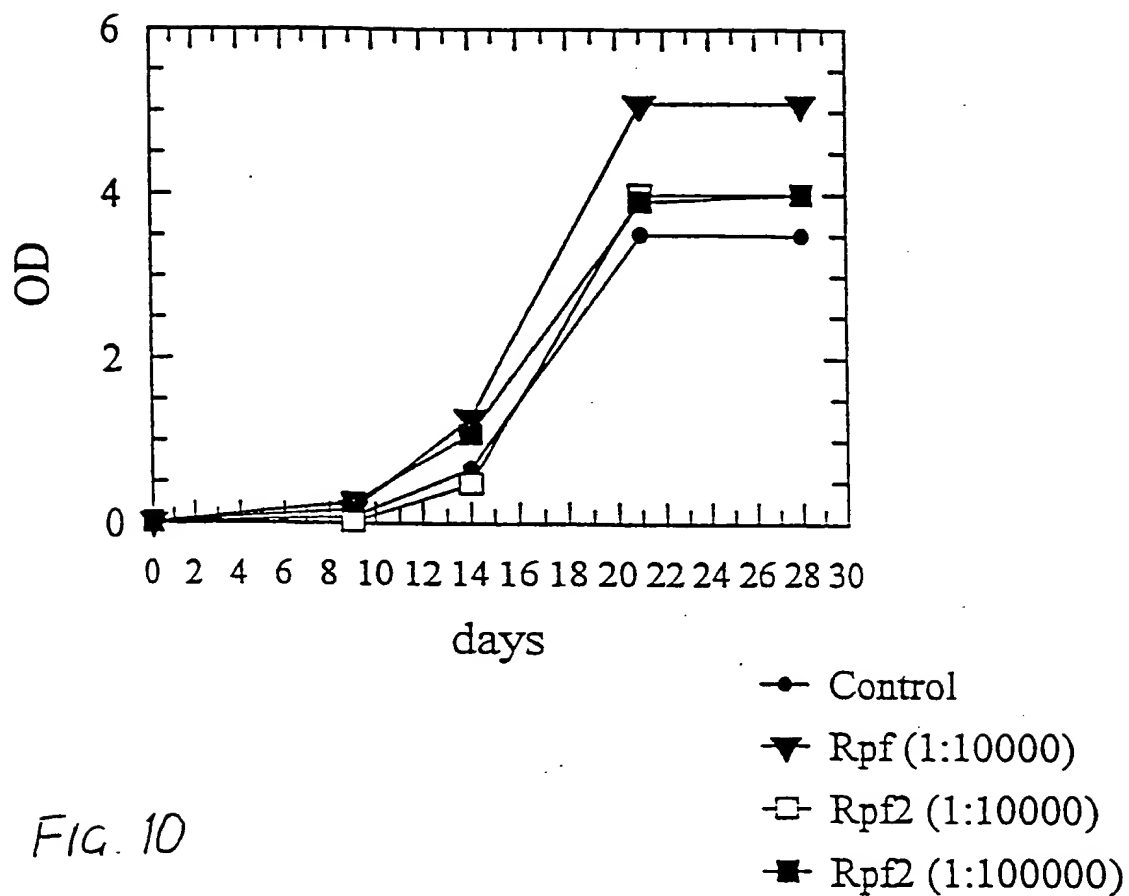


FIG. 10